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Raw Sequence Listing
Patent Application US/07/661,070

04/22/91
09:27:22

SEQUENCE LISTING

(1) GENERAL INFORMATION:

(i) APPLICANT: Huston, James S
Charette, Marc F
Cohen, Charles M
Crea, Roberto
Keck, Peter C
Oppermann, Hermann
Rueger, David C
Ridge, Richard J

(ii) TITLE OF INVENTION: Product and Process for the Production,
Isolation and Purification of Recombinant Polypeptides

(iii) NUMBER OF SEQUENCES: 14

(iv) CORRESPONDENCE ADDRESS:

(A) ADDRESSEE: Creative BioMolecules
(B) STREET: 35 South Street
(C) CITY: Hopkinton
(D) STATE: MA
(E) COUNTRY: USA
(F) ZIP: 01748

(v) COMPUTER READABLE FORM:

(A) MEDIUM TYPE: Floppy disk
(B) COMPUTER: IBM PC compatible
(C) OPERATING SYSTEM: PC-DOS/MS-DOS
(D) SOFTWARE: PatentIn Release #1.0, Version #1.25

(vi) CURRENT APPLICATION DATA:

(A) APPLICATION NUMBER: US 07/661,070
(B) FILING DATE: 26-FEB-1991
(C) CLASSIFICATION: 435/68
536/27
530/300
530/350

(viii) ATTORNEY/AGENT INFORMATION:

(A) NAME: Lunn, Paul G.
(B) REGISTRATION NUMBER: 32,743
(C) REFERENCE/DOCKET NUMBER: CRP-008DV

(ix) TELECOMMUNICATION INFORMATION:

(A) TELEPHONE: (508) 435-9001
(B) TELEFAX: (508) 435-6951

(2) INFORMATION FOR SEQ ID NO:1:

Patent Application US/07/661,070

54 (i) SEQUENCE CHARACTERISTICS:
55 (A) LENGTH: 4 amino acids
56 (B) TYPE: amino acid
57 (C) STRANDEDNESS: single
58 (D) TOPOLOGY: linear
59
60 (ii) MOLECULE TYPE: peptide
61
62 (iii) HYPOTHETICAL: NO
63
64 (iv) ANTI-SENSE: NO
65
66 (v) FRAGMENT TYPE: internal
67
68
69
70 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:
71
72 Ile Glu Gly Arg
73 1
74
75 (2) INFORMATION FOR SEQ ID NO:2:
76
77 (i) SEQUENCE CHARACTERISTICS:
78 (A) LENGTH: 21 base pairs
79 (B) TYPE: nucleic acid
80 (C) STRANDEDNESS: double
81 (D) TOPOLOGY: linear
82
83 (ii) MOLECULE TYPE: cDNA
84
85 (iii) HYPOTHETICAL: NO
86
87 (iv) ANTI-SENSE: NO
88
89 (v) FRAGMENT TYPE: N-terminal
90
91
92 (ix) FEATURE:
93 (A) NAME/KEY: CDS
94 (B) LOCATION: 1..21
95
96
97 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:
98
99 GCT AAA AAC CTT AAC GAA GCT
100 Ala Lys Asn Leu Asn Glu Ala
101 1 5
102
103
104 (2) INFORMATION FOR SEQ ID NO:3:
105
106 (i) SEQUENCE CHARACTERISTICS:

Patent Application US/07/661,070

107 (A) LENGTH: 7 amino acids
108 (B) TYPE: amino acid
109 (D) TOPOLOGY: linear
110
111 (ii) MOLECULE TYPE: protein
112
113 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:
114
115 Ala Lys Asn Leu Asn Glu Ala
116 1 5
117
118 (2) INFORMATION FOR SEQ ID NO:4:
119
120 (i) SEQUENCE CHARACTERISTICS:
121 (A) LENGTH: 13 amino acids
122 (B) TYPE: amino acid
123 (C) STRANDEDNESS: single
124 (D) TOPOLOGY: linear
125
126 (ii) MOLECULE TYPE: peptide
127
128 (iii) HYPOTHETICAL: NO
129
130 (iv) ANTI-SENSE: NO
131
132 (v) FRAGMENT TYPE: internal
133
134
135
136 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:
137
138 Met Lys Ala Ile Phe Val Leu Lys Gly Ser Leu Asp Glu
139 1 5 10
140
141 (2) INFORMATION FOR SEQ ID NO:5:
142
143 (i) SEQUENCE CHARACTERISTICS:
144 (A) LENGTH: 16 amino acids
145 (B) TYPE: amino acid
146 (C) STRANDEDNESS: single
147 (D) TOPOLOGY: linear
148
149 (ii) MOLECULE TYPE: peptide
150
151 (iii) HYPOTHETICAL: NO
152
153 (iv) ANTI-SENSE: NO
154
155 (v) FRAGMENT TYPE: internal
156
157
158
159 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:

Patent Application US/07/661,070

160
161 Met Lys Ala Ile Phe Val Leu Lys Gly Ser Leu Asp Arg Asp Leu Glu
162 1 5 10 15
163
164

165 (2) INFORMATION FOR SEQ ID NO:6:

- 166
167 (i) SEQUENCE CHARACTERISTICS:
168 (A) LENGTH: 59 amino acids
169 (B) TYPE: amino acid
170 (C) STRANDEDNESS: single
171 (D) TOPOLOGY: linear
172

173 (ii) MOLECULE TYPE: protein
174

175 (iii) HYPOTHETICAL: NO
176

177 (iv) ANTI-SENSE: NO
178

179 (v) FRAGMENT TYPE: N-terminal
180
181
182

183 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:

184
185 Met Lys Ala Ile Phe Val Leu Lys Gly Ser Leu Asp Arg Asp Leu Asp
186 1 5 10 15
187

188 Ser Arg Leu Asp Leu Asp Val Arg Thr Asp His Lys Asp Leu Ser Asp
189 20 25 30
190

191 His Leu Val Leu Val Asp Leu Ala Arg Asn Asp Leu Ala Arg Ile Val
192 35 40 45
193

194 Thr Pro Gly Ser Arg Tyr Val Ala Asp Leu Glu
195 50 55
196

197 (2) INFORMATION FOR SEQ ID NO:7:

- 198
199 (i) SEQUENCE CHARACTERISTICS:
200 (A) LENGTH: 4 amino acids
201 (B) TYPE: amino acid
202 (C) STRANDEDNESS: single
203 (D) TOPOLOGY: linear
204

205 (ii) MOLECULE TYPE: peptide
206

207 (iii) HYPOTHETICAL: NO
208

209 (iv) ANTI-SENSE: NO
210

211 (v) FRAGMENT TYPE: internal
212

Patent Application US/07/661,070

213
214
215 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:
216
217 Glu Phe Met Arg
218 1
219
220 (2) INFORMATION FOR SEQ ID NO:8:
221
222 (i) SEQUENCE CHARACTERISTICS:
223 (A) LENGTH: 10 amino acids
224 (B) TYPE: amino acid
225 (C) STRANDEDNESS: single
226 (D) TOPOLOGY: linear
227
228 (ii) MOLECULE TYPE: peptide
229
230 (iii) HYPOTHETICAL: NO
231
232 (iv) ANTI-SENSE: NO
233
234 (v) FRAGMENT TYPE: internal
235
236
237
238 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:
239
240 Glu Phe Asp Pro Pro Pro Lys Phe Met Arg
241 1 5 10
242
243 (2) INFORMATION FOR SEQ ID NO:9:
244
245 (i) SEQUENCE CHARACTERISTICS:
246 (A) LENGTH: 13 amino acids
247 (B) TYPE: amino acid
248 (C) STRANDEDNESS: single
249 (D) TOPOLOGY: linear
250
251 (ii) MOLECULE TYPE: peptide
252
253 (iii) HYPOTHETICAL: NO
254
255 (iv) ANTI-SENSE: NO
256
257 (v) FRAGMENT TYPE: internal
258
259
260
261 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:
262
263 Glu Phe Asp Pro Pro Pro Met Pro Arg Lys Phe Met Arg
264 1 5 10
265

Patent Application US/07/661,070

266 (2) INFORMATION FOR SEQ ID NO:10:

267

268 (i) SEQUENCE CHARACTERISTICS:

269 (A) LENGTH: 20 amino acids

270 (B) TYPE: amino acid

271 (C) STRANDEDNESS: single

272 (D) TOPOLOGY: linear

273

274 (ii) MOLECULE TYPE: peptide

275

276 (iii) HYPOTHETICAL: NO

277

278 (iv) ANTI-SENSE: NO

279

280 (v) FRAGMENT TYPE: internal

281

282

283

284 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:

285

286 Glu Phe Asp Pro Pro Pro Met Pro Arg Met Pro Asp Pro Glu Leu Arg

287 1 5 10 15

288

289 Lys Phe Met Arg

290 20

291

292 (2) INFORMATION FOR SEQ ID NO:11:

293

294 (i) SEQUENCE CHARACTERISTICS:

295 (A) LENGTH: 193 amino acids

296 (B) TYPE: amino acid

297 (C) STRANDEDNESS: single

298 (D) TOPOLOGY: linear

299

300 (ii) MOLECULE TYPE: protein

301

302 (iii) HYPOTHETICAL: NO

303

304 (iv) ANTI-SENSE: NO

305

306 (v) FRAGMENT TYPE: N-terminal

307

308

309

310 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:

311

312 Met Lys Ala Ile Phe Val Leu Lys Gly Ser Leu Asp Arg Asp Leu Asp

313 1 5 10 15

314

315 Ser Arg Ile Glu Leu Glu Met Arg Thr Asp His Lys Glu Leu Ser Glu

316 20 25 30

317

318 His Leu Met Leu Val Asp Leu Ala Arg Asn Asp Leu Ala Arg Ile Cys

Raw Sequence Listing
Patent Application US/07/661,070

04/22/91
09:27:33

	35	40	45
319			
320			
321	Thr Pro Gly Ser Arg Tyr Val Ala Asp Leu Thr Lys Val Asp Arg Tyr		
322	50	55	60
323			
324	Ser Tyr Val Met His Leu Val Ser Arg Val Val Gly Glu Leu Arg His		
325	65	70	75
326			
327	Asp Leu Asp Ala Leu His Ala Tyr Arg Ala Cys Met Asn Met Gly Thr		
328		85	90
329			95
330	Leu Ser Gly Ala Pro Lys Val Arg Ala Met Gln Leu Ile Ala Glu Ala		
331		100	105
332			110
333	Glu Gly Arg Arg Arg Gly Ser Tyr Gly Gly Ala Val Gly Tyr Phe Thr		
334		115	120
335			125
336	Ala His Gly Asp Leu Asp Thr Cys Ile Val Ile Arg Ser Ala Leu Val		
337		130	135
338			140
339	Glu Asn Gly Ile Ala Thr Val Gln Ala Gly Ala Gly Val Val Leu Asp		
340		145	150
341			155
342	Ser Val Pro Gln Ser Glu Ala Asp Glu Thr Arg Asn Lys Ala Arg Ala		
343		165	170
344			175
345	Val Leu Arg Ala Ile Ala Thr Ala His His Ala Gln Glu Phe Pro Gly		
346		180	185
347			190
348	Glu		
349			
350			

(2) INFORMATION FOR SEQ ID NO:12:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 59 amino acids
- (B) TYPE: amino acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: protein

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(v) FRAGMENT TYPE: N-terminal

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:12:

Met Lys Ala Ile Phe Val Leu Lys Gly Ser Leu Asp Arg Asp Leu Asp

Patent Application US/07/661,070

372 1 5 10 15
373
374 Ser Arg Leu Asp Leu Asp Val Arg Thr Asp His Lys Asp Leu Ser Asp
375 20 25 30
376
377 His Leu Val Leu Val Asp Leu Ala Arg Asn Asp Leu Ala Arg Ile Val
378 35 40 45
379
380 Thr Pro Gly Ser Arg Tyr Val Ala Asp Leu Glu
381 50 55
382

(2) INFORMATION FOR SEQ ID NO:13:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 21 amino acids
- (B) TYPE: amino acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(v) FRAGMENT TYPE: internal

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:13:

403 Met Lys Ala Ile Phe Val Leu Lys Gly Ser Leu Asp Arg Asp Leu Glu
404 1 5 10 15
405
406 Phe Met Pro Pro Cys
407 20
408

(2) INFORMATION FOR SEQ ID NO:14:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 19 amino acids
- (B) TYPE: amino acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(v) FRAGMENT TYPE: internal

Raw Sequence Listing
Patent Application US/07/661,070

04/22/91
09:27:53

425
426
427
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433

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:14:

Met	Lys	Ala	Ile	Phe	Val	Leu	Lys	Gly	Ser	Leu	Asp	Arg	Asp	Leu	Glu
1				5				10					15		
Phe Met Cys															

PAGE: 1

SEQUENCE VERIFICATION REPORT
PATENT APPLICATION US/07/661,070

DATE: 04/22/91
TIME: 09:27:54

LINE ERROR

ORIGINAL TEXT

35 Wrong application Serial Number
36 Wrong Filing Date
37 Wrong Classification

(A) APPLICATION NUMBER: US 07/661,070
(B) FILING DATE: 26-FEB-1991
(C) CLASSIFICATION: 435/68

PAGE: 1

SEQUENCE MISSING ITEM REPORT
PATENT APPLICATION US/07/661,070

DATE: 04/22/91
TIME: 09:27:54

MANDATORY IDENTIFIER THAT WAS NOT FOUND

PRIOR APPLICATION DATA
APPLICATION NUMBER
FILING DATE

PAGE: 1

SEQUENCE CORRECTION REPORT
PATENT APPLICATION US/07/661,070

DATE: 04/22/91
TIME: 09:27:54

LINE ORIGINAL TEXT

CORRECTED TEXT